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                                                                                 300
Ħ
                                                                                 360
Ž.
      agaaaaccat ctccaaagcc aaagggcagc cccgagaacc acaggtgtac accctgcccc
                                                                                 420
l
Heli
      catcccggga tgagctgacc aagaaccagg tcagcctgac ctgcctggtc aaaggcttct
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Lil
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Į.i.
                                                                                 733
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       <223> Synthetic sequence with 4 tandem copies of the GAS binding site
              found in the IRF1 promoter (Rothman et al., Immunity 1:457-468
              (1994)), 18 nucleotides complementary to the SV40 early promoter,
             and a Xho I restriction site.
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                                                                       180
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                                                                       300
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                                                                       420
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acctccagec tcacaccatc tcttcagect agcaagttgc tggagggagt ctataaccta
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ccaggagcca gccagccatt tgtatcaaga aatagaaatc tgccagggta cagtggctca
                                                                      300
cacctataat cccagcgctt tgggaggcta agttctagga caaggcaaga agaaaqcaaq
                                                                      360
aagctgtaaa tcccattcct ctgggtctca atttcaccct cagttcaagg agctgagtag
                                                                      420
gcagaggcaa aggctatact caacacacgt gcaattgaaa gcaggcgagg caaaaccagg
                                                                      480
gcagaggaaa ggaaaggggt gtgtgtgtag aactgctcag ggtagactgg agacaaaagc
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                                                                      629
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gtagaaaaga tactcatcca ctgtgggttt tggtttcgcc gtcaccccac tgcctcactg
                                                                      180
                                                                      240
gattgtgagg atcatatgcg acaatgtatt tgaaaacgac tagaacatta tcggaggaag
gtggactctg aagtagtcgc tgtagactat ggatgtagaa caagggtttg gagcccttcg
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gacatggttc taacgcggcc tgacttcttg ctggctacat gaccttggac tacataatca
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cgcctcttaa atgggaggtg atgacagcta tccttgagga ccttagagag aactgatttc
                                                                      420
ttagtaccca gcctcacaaa tagtgcatca cttcatggag ttatgttggg ataaatgtgt
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ggagaagcca gggaatcgcc tagactctcg cactgaaaat tgtctctcca gctgtgtaga
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cogetteatt gacaccacte ttgccateae cagteggttt geccagattg tacagaaagg
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agggcagtag gccatccccc aggagaatga cagaagcaaa ggacttgtta ctaagcagat
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ttaagggtca gtgggggaag gctatcaacc cattgtcaga tcagcatcag gctgttatca
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<222> (838)..(838)
<223> n equals a,t,g, or c
<221> misc_feature
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gctgatctcc tgcctgatgt ttctgtctca gagccaaggc caagaggccc agacagagtt
                                                                      180
gccccaggcc cggatcagct gcccagaagg caccaatgcc tatcgctcct actgctacta
                                                                      240
                                                                      300
ctttaatgaa gaccgtgaga cctgggttga tgcagatctc tattgccaga acatgaattc
aggeaacctq qtgtctqtgc tcacccaggc cgagggtgcc tttgtggcct cactgattaa
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                                                                      420
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                                                                      660
gaccatetet ecaacteaac teaacetgga cactetette tetgetgagt ttgeettgtt
                                                                      720
                                                                      780
aatcttcaat agttttacct accccagtct ttggaaccyt aaataataaa aataaacatg
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<223> n equals a,t,g, or c
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ggcccagggc cctgcgggcc accccccac catggcccag ggccctgcgg gccaccccc
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caccatggtc cagggccctg cgggccaccc cctggccatg gcccagggcc ctgcgggcca
                                                                      240
                                                                      300
ccccccacc atggtccagg gccctgcggg cctcccctg gccatggccc aggtcaccca
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ttgcatgcaa tgactatttt gtagtacaca tgaagcagaa gggaaagaag taggcagaaa
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tgagcagttc gctcctccct gataagagtt gtcccaaagg gtcgcttaag gaatctgccc
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                                                                      300
cacagcttcc cccatagaag gatttcatga gcagatcagg acacttagca aatgtaaaaa
taaaatctaa ctctcatttg acaagcagag aaagaaaagt taaataccag ataagctttt
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<211> 1637
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<213> Homo sapiens
<220>
<221> misc_feature
<222> (738)..(738)
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<222> (771)..(771)
<223> n equals a,t,q, or c
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cgtggacaat ggctacttgg agggactggt gcgcggcctg aaggccgggg tgctcagcca
                                                                     180
ggccgactac ctcaacctgg tgcagtgcga gacgctagag gacttgaaac tgcatctgca
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gagcactgat tatggtaact tcctggccaa cgaggcatca cctctgacgg tgtcagtcat
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cgatgaccgg ctcaaggaga agatggtggt ggagttccgc cacatgagga accatgccta
                                                                     360
tgagccactc gccagcttcc tagacttcat tacttacagt tacatgatcg acaacgtgat
                                                                     420
cctgctcatc acaggcacgc tgcaccagcg ctccatcgct gagctcgtgc ccaagtqcca
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cccactaggc agcttcgagc agatggaggc cgtgaacatt gctcagacac ctgctgagct
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gcaggacett gacgagatga acategagat cateegcaac accetetaca aggeetacet
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tegtgtggat egetgaatgt ategeeeage gecacegege caaaategae aactacatee
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ctatetteta gegteetgge ceaaggetet caattgeact etttgtgtgt gtgtgtgt
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gtgtgcgcgt gtgtgtgcgt gtgtgtgtat gtggtctgtg acaagcctgt ggctcacctg
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cctgtccggg gtgtagtacg ctgtcctagc ggctgcccag ttctcctgac cctcttagag
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geatgteact tteatgttee teectaacte cetgacetga gaaceetggg geetggggge
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1620
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<211> 1471
<212> DNA
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                                                                     120
ctgtgtcatg gtggttttgg ttctaaacag gtatgcagaa ggtccccgtt acactttcca
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ataatgaaaa atgtttataa ttctaaatac agcaacccat gtaagacatg ttcatgtatc
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                                                                    420
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                                                                    480
aaaaagagac aggaaaatta aaataatcaa atctatataa atacatgaat catgctgaca
                                                                    540
acacaggact gatttttcgt ttgattattt taacacagac agatgtaaat cccaaaagac
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                                                                   1320
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acticitytat gtccggcaga ggaacgcttg gcatccaatc tgatgtatcc cagtgatacc
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<210> 29
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<212> DNA
<213> Homo sapiens
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tgctggtcgt cgcccactgc tgctgctgca gctcccccgg gccccgcagg gaaagcccca
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420
aa
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<211> 166
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<213> Homo sapiens
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Arg Ile Ser Cys Pro Glu Gly Thr Asn Ala Tyr Arg Ser Tyr Cys Tyr
Tyr Phe Asn Glu Asp Arg Glu Thr Trp Val Asp Ala Asp Leu Tyr Cys
Gln Asn Met Asn Ser Gly Asn Leu Val Ser Val Leu Thr Gln Ala Glu
Gly Ala Phe Val Ala Ser Leu Ile Lys Glu Ser Gly Thr Asp Asp Phe
Asn Val Trp Ile Gly Leu His Asp Pro Lys Lys Asn Arg Arg Trp His
                               105
Trp Ser Ser Gly Ser Leu Val Ser Tyr Lys Ser Trp Gly Ile Gly Ala
Pro Ser Ser Val Asn Pro Gly Tyr Cys Val Ser Leu Thr Ser Ser Thr
Gly Phe Gln Lys Trp Lys Asp Val Pro Cys Glu Asp Lys Phe Ser Phe
                   150
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Val Cys Lys Phe Lys Asn
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 <211> 93
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 <213> Homo sapiens
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 Pro Pro Gly His Cys Gly Pro Pro Pro Gly His Gly Pro Gly Pro Cys 20 25 30
 Gly Pro Pro Pro His His Gly Pro Gly Pro Cys Gly Pro Pro Pro His
 His Gly Pro Gly Pro Cys Gly Pro Pro Pro Gly His Gly Pro Gly Pro 50 60
Cys Gly Pro Pro Pro His His Gly Pro Gly Pro Cys Gly Pro Pro 65 70 75 80
Gly His Gly Pro Gly His Pro Pro Pro Gly Pro His His
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<211> 97
<212> PRT
<213> Homo sapiens
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His Lys Phe Ala Gly Asp Lys Gly Tyr Leu Thr Lys Glu Asp Leu Arg 20 25 30
Val Leu Met Glu Lys Glu Phe Pro Gly Phe Leu Glu Asn Gln Lys Asp
Pro Leu Ala Val Asp Lys Ile Met Lys Asp Leu Asp Glr Cys Arg Asp
                           55
Gly Lys Val Gly Phe Gln Ser Phe Phe Ser Leu Ile Ala Gly Leu Thr
Ile Ala Cys Asn Asp Tyr Phe Val Val His Met Lys Gln Lys Gly Lys 85 90 95
Lys
<210> 33
<211> 351
<212> PRT
<213> Homo sapiens
<400> 33
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Met Ser Phe Phe Pro Glu Leu Tyr Phe Asn Val Asp Asn Gly Tyr Leu  $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$ 

Glu Gly Leu Val Arg Gly Leu Lys Ala Gly Val Leu Ser Gln Ala Asp Tyr Leu Asn Leu Val Gln Cys Glu Thr Leu Glu Asp Leu Lys Leu His Leu Gln Ser Thr Asp Tyr Gly Asn Phe Leu Ala Asn Glu Ala Ser Pro Leu Thr Val Ser Val Ile Asp Asp Arg Leu Lys Glu Lys Met Val Val 65 70 75 80 Glu Phe Arg His Met Arg Asn His Ala Tyr Glu Pro Leu Ala Ser Phe Leu Asp Phe Ile Thr Tyr Ser Tyr Met Ile Asp Asn Val Ile Leu Leu Ile Thr Gly Thr Leu His Gln Arg Ser Ile Ala Glu Leu Val Pro Lys Cys His Pro Leu Gly Ser Phe Glu Gln Met Glu Ala Val Asn Ile Ala Gln Thr Pro Ala Glu Leu Tyr Asn Ala Ile Leu Val Asp Thr Pro Leu Ala Ala Phe Phe Gln Asp Cys Ile Ser Glu Gln Asp Leu Asp Glu Met Asn Ile Glu Ile Ile Arg Asn Thr Leu Tyr Lys Ala Tyr Leu Glu Ser Phe Tyr Lys Phe Cys Thr Leu Leu Gly Gly Thr Thr Ala Asp Ala Met Cys Pro Ile Leu Glu Phe Glu Ala Asp Arg Arg Ala Phe Ile Ile Thr Ile Asn Ser Phe Gly Thr Glu Leu Ser Lys Glu Asp Arg Ala Lys Leu 225 230 235 Phe Pro His Cys Gly Arg Leu Tyr Pro Glu Gly Leu Ala Gln Leu Ala Arg Ala Asp Asp Tyr Glu Gln Val Lys Asn Val Ala Asp Tyr Tyr Pro 260 265 270 Glu Tyr Lys Leu Leu Phe Glu Gly Ala Gly Ser Asn Pro Gly Asp Lys
275
280 Thr Leu Glu Asp Arg Phe Phe Glu His Glu Val Lys Leu Asn Lys Leu Ala Phe Leu Asn Gln Phe His Phe Gly Val Phe Tyr Ala Phe Val Lys Leu Lys Glu Gln Glu Cys Arg Asn Ile Val Trp Ile Ala Glu Cys Ile 330 Ala Gln Arg His Arg Ala Lys Ile Asp Asn Tyr Ile Pro Ile Phe 340

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<211> 127
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<212> PRT

<213> Homo sapiens

<400> 34

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Ala Phe Met Lys Ala Ile Gly Leu Pro Glu Glu Leu Ile Gln Lys Gly 20 25 30

Lys Asp Ile Lys Gly Val Ser Glu Ile Val Gln Asn Gly Lys His Phe 35 40 45

Lys Phe Thr Ile Thr Ala Gly Ser Lys Val Ile Gln Asn Glu Phe Thr 50 60

Val Gly Glu Glu Cys Glu Leu Glu Thr Met Thr Gly Glu Lys Val Lys 65 70 75 80

Thr Val Val Gln Leu Glu Gly Asp Asn Lys Leu Val Thr Thr Phe Lys 85 90 95

Asn Ile Lys Ser Val Thr Glu Leu Asn Gly Asp Ile Ile Thr Asn Thr 100 105 110

Met Thr Leu Gly Asp Ile Val Phe Lys Arg Ile Ser Lys Arg Ile 115 120 125

<210> 35

<211> 219

<212> PRT

<213> Homo sapiens

<400> 35

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Asn Tyr Glu Val Ile Asp Glu Gln Thr Pro Leu Tyr Ser Ala Asp Pro 35 40 45

Asn Ala Ile Asp Thr Asp Tyr Tyr Pro Gly Gly Tyr Asp Ile Glu Ser 50 60

Asp Phe Pro Pro Pro Pro Glu Asp Phe Pro Ala Ala Asp Glu Leu Pro 65 70 75 80

Pro Leu Pro Pro Glu Phe Ser Asn Gln Phe Glu Ser Ile His Pro Pro 85 90 95

Arg Asp Met Pro Ala Ala Gly Ser Leu Gly Ser Ser Ser Arg Asn Arg 100 105 110

Gln Arg Phe Asn Leu Asn Gln Tyr Leu Pro Asn Phe Tyr Pro Leu Asp 115 120 125

Met Ser Glu Pro Gln Thr Lys Gly Thr Gly Glu Asn Ser Thr Cys Arg 130 135 140

Glu Pro His Ala Pro Tyr Pro Pro Gly Tyr Gln Arg His Phe Glu Ala

<210> 36 <211> 256 <212> PRT <213> Homo sapiens

Ala Ala Asp Asp Phe Arg Thr Lys Phe Glu Thr Glu Gln Ala Leu Arg 20 25 30

Met Ser Val Glu Ala Asp Ile Asn Gly Leu Arg Arg Val Leu Asp Glu 35 40 45

Leu Thr Leu Ala Arg Thr Asp Leu Glu Met Gln Ile Glu Gly Leu Lys
50 60

Glu Glu Leu Ala Tyr Leu Lys Lys Asn His Glu Glu Glu Ile Ser Thr 65 70 75 80

Leu Arg Gly Gln Val Gly Gly Gln Val Ser Val Glu Val Asp Ser Ala 85 90 95

Pro Gly Thr Asp Leu Ala Lys Ile Leu Ser Asp Met Arg Ser Gln Tyr 100 105 110

Glu Val Met Ala Glu Gln Asn Arg Lys Asp Ala Glu Ala Trp Phe Thr 115 120 125

Ser Arg Thr Glu Glu Leu Asn Arg Glu Val Ala Gly His Thr Glu Gln 130 135

Leu Gln Met Ser Arg Ser Glu Val Thr Asp Leu Arg Arg Thr Leu Gln 145 150 155 160

Gly Leu Glu Ile Glu Leu Gln Ser Gln Leu Ser Met Lys Ala Ala Leu 165 170 175

Glu Asp Thr Leu Ala Glu Thr Glu Ala Arg Phe Gly Ala Gln Leu Ala 180 185 190

His Ile Gln Ala Leu Ile Ser Gly Ile Glu Ala Gln Leu Gly Asp Val 195 200 205

Arg Ala Asp Ser Glu Arg Gln Asn Gln Glu Tyr Gln Arg Leu Met Asp 210 215 220

Ile Lys Ser Arg Leu Glu Gln Glu Ile Ala Thr Tyr Arg Ser Leu Leu 225 230 235 240 Glu Gly Gln Glu Asp His Tyr Asn Asn Leu Ser Ala Ser Lys Val Leu 250

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<210> 37
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<400> 37

Met Ser Val Ser Glu Leu Lys Ala Gln Ile Thr Gln Lys Ile Gly Val

His Ala Phe Gln Gln Arg Leu Ala Val His Pro Ser Gly Val Ala Leu

Gln Asp Arg Val Pro Leu Ala Ser Gln Gly Leu Gly Pro Gly Ser Thr

Val Leu Leu Val Val Asp Lys Cys Asp Glu Pro Leu Ser Ile Leu Val 50 60

Arg Asn Asn Lys Gly Arg Ser Ser Thr Tyr Glu Val Arg Leu Thr Gln 65 70 75 80

Thr Val Ala His Leu Lys Gln Gln Val Ser Gly Leu Glu Gly Val Gln

Asp Asp Leu Phe Trp Leu Thr Phe Glu Gly Lys Pro Leu Glu Asp Gln

Leu Pro Leu Gly Glu Tyr Gly Leu Lys Pro Leu Ser Thr Val Phe Met 115 120 125

Asn Leu Arg Leu Arg Gly Gly Gly Thr Glu Pro Gly Gly Arg Ser 130 135 140

<210> 38

<211> 41

<212> PRT

<213> Homo sapiens

<400> 38

Met Thr Gly Ala Gly Leu Gly Arg Asp Ser Gly Arg Trp Arg Glu Val

Ser Phe Phe Gly Glu Thr Glu Arg Ala Arg Gly Gly Thr Val Gly Arg

Gly Arg Thr Arg Leu Arg Arg Gln Glu

<sup>&</sup>lt;211> 143

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

<sup>&</sup>lt;210> 39 <211> 118

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

<400> 39 Met Gln Thr Pro Pro Cys Phe Leu His Leu Ala Asp Tyr Leu Tyr Pro Glu Gln Leu Lys Met Thr Val Val Lys Leu Ile Ser His Arg Glu Cys 20 25 30Gln Gln Pro His Tyr Tyr Gly Ser Glu Val Thr Thr Lys Met Leu Cys Ala Ala Asp Pro Gln Trp Lys Thr Asp Ser Cys Gln Gly Asp Ser Gly 50 60 Gly Pro Leu Val Cys Ser Leu Gln Gly Arg Met Thr Leu Thr Gly Ile 65 70 75 80 Val Ser Trp Gly Arg Gly Cys Ala Leu Lys Asp Lys Pro Gly Val Tyr Thr Arg Val Ser His Phe Leu Pro Trp Ile Arg Ser His Thr Lys Glu 100 105 110Glu Asn Gly Leu Ala Leu

<210> 40 <211> 110 <212> PRT <213> Homo sapiens

115

<400> 40

Pro Arg Val Arg Pro Glu Ala Gly Ser Ser Leu Cys Ser Pro Gly Pro 1 5 10 15 Ala Trp Leu Gly Glu Leu Glu Ala Ser Arg Trp His Gly Ala Arg 20 25 30

Gln Asp Gly Cys Val His Arg Gly Ala Gly Gly His Gly Ser Gly Ser 35 40 45

Pro Gly Glu Thr Glu Glu Pro Pro Val Phe Pro Val His Met Gly His  $50 \hspace{1.5cm} 55 \hspace{1.5cm} 60 \hspace{1.5cm}$ 

Cys Cys Leu His Cys Phe Pro His Leu His Gly His Arg Ala Ala Pro 65 70 75 80

Ala Ala Gly Arg Arg Pro Leu Leu Leu Gln Leu Pro Arg Ala 85 90 95

Pro Gln Gly Lys Pro Gln Glu Gly Lys Thr Gln Gly Ser Gly

<210> 41 <211> 63 <212> PRT

<213> Homo sapiens

<400> 41 Asp Thr Thr Thr Arg Asp Phe Thr Gln Leu Asn Glu Leu Gln Cys Arg

Phe Pro Arg Arg Leu Val Val Leu Gly Phe Pro Cys Asn Gln Phe Gly

W

1.5

20 25 30

His Gln Ser Arg Arg Asp Arg Ser Ser Lys Pro Ser Phe Glu Met Ser 35 40 45

Leu Gln Pro Gln Lys Tyr Leu Gln Pro His Thr Ile Ser Ser Ala 50 60

<210> 42

<211> 61

<212> PRT

<213> Homo sapiens

<400> 42

Tyr Pro Ala Ser Gln Ile Val His His Phe Met Glu Leu Cys Trp Asp 1 10 15

Lys Cys Val Glu Lys Pro Gly Asn Arg Leu Asp Ser Arg Thr Glu Asn 20 25 30

Cys Leu Ser Ser Cys Val Asp Arg Phe Ile Asp Thr Thr Leu Ala Ile  $35 \hspace{1cm} 40 \hspace{1cm} 45$ 

Thr Ser Arg Phe Ala Gln Ile Val Gln Lys Gly Gln 50 60

<210> 43

<211> 166

<212> PRT

<213> Homo sapiens

<400> 43

Met Ala Gln Thr Ser Ser Tyr Phe Met Leu Ile Ser Cys Leu Met Phe 1 15

Leu Ser Gln Ser Gln Gly Gln Glu Ala Gln Thr Glu Leu Pro Gln Ala

Arg Ile Ser Cys Pro Glu Gly Thr Asn Ala Tyr Arg Ser Tyr Cys Tyr

Tyr Phe Asn Glu Asp Arg Glu Thr Trp Val Asp Ala Asp Leu Tyr Cys
50 60

Gln Asn Met Asn Ser Gly Asn Leu Val Ser Val Leu Thr Gln Ala Glu 65 70 75 80

Gly Ala Phe Val Ala Ser Leu Ile Lys Glu Ser Gly Thr Asp Asp Phe \$90\$

Asn Val Trp Ile Gly Leu His Asp Pro Lys Lys Asn Arg Arg Trp His
100 105 110

Trp Ser Ser Gly Ser Leu Val Ser Tyr Lys Ser Trp Gly Ile Gly Ala 115 120 125

Pro Ser Ser Val Asn Pro Gly Tyr Cys Val Ser Leu Thr Ser Ser Thr 130 135 140

Gly Phe Gln Lys Trp Lys Asp Val Pro Cys Glu Asp Lys Phe Ser Phe 145 150 155 160

<220> <221> SITE <222> (216)

Val Cys Lys Phe Lys Asn 165

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<210> 44
<211> 93
<212> PRT
<213> Homo sapiens
<400> 44
Met Asp Pro Gly Pro Lys Gly His Cys His Cys Gly Gly His Gly His
Pro Pro Gly His Cys Gly Pro Pro Pro Gly His Gly Pro Gly Pro Cys
Gly Pro Pro Pro His His Gly Pro Gly Pro Cys Gly Pro Pro Pro His
His Gly Pro Gly Pro Cys Gly Pro Pro Pro Gly His Gly Pro Gly Pro
Cys Gly Pro Pro Pro His His Gly Pro Gly Pro Cys Gly Pro Pro Pro 65 70 75 80
Gly His Gly Pro Gly His Pro Pro Pro Gly Pro His His
                 85
<210> 45
<211> 56
<212> PRT
<213> Homo sapiens
<400> 45
His Glu Glu Asn Gln Lys Asp Pro Leu Ala Val Asp Lys Ile Met Lys
Asp Leu Asp Gln Cys Arg Asp Gly Lys Val Gly Phe Gln Ser Phe Phe
Ser Leu Ile Ala Gly Leu Thr Ile Ala Cys Asn Asp Tyr Phe Val Val
His Met Lys Gln Lys Gly Lys Lys
<210> 46
<211> 239
<212> PRT
<213> Homo sapiens
<220>
<221> SITE
<222> (215)
<223> Xaa equals any of the naturally occurring L-amino acids
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<223> Xaa equals any of the naturally occurring L-amino acids

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<220>
<221> SITE
<222> (227)
<223> Xaa equals any of the naturally occurring L-amino acids
<220>
<221> SITE
<222> (236)
<223> Xaa equals any of the naturally occurring L-amino acids
Met Ser Phe Phe Pro Glu Leu Tyr Phe Asn Val Asp Asn Gly Tyr Leu
Glu Gly Leu Val Arg Gly Leu Lys Ala Gly Val Leu Ser Gln Ala Asp
Tyr Leu Asn Leu Val Gln Cys Glu Thr Leu Glu Asp Leu Lys Leu His
Leu Gln Ser Thr Asp Tyr Gly Asn Phe Leu Ala Asn Glu Ala Ser Pro
Leu Thr Val Ser Val Ile Asp Asp Arg Leu Lys Glu Lys Met Val Val
Glu Phe Arg His Met Arg Asn His Ala Tyr Glu Pro Leu Ala Ser Phe
Leu Asp Phe Ile Thr Tyr Ser Tyr Met Ile Asp Asn Val Ile Leu Leu
                                 105
Ile Thr Gly Thr Leu His Gln Arg Ser Ile Ala Glu Leu Val Pro Lys 115 120 125
Cys His Pro Leu Gly Ser Phe Glu Gln Met Glu Ala Val Asn Ile Ala
                         135
Gln Thr Pro Ala Glu Leu Tyr Asn Ala Ile Leu Val Asp Thr Pro Leu
                                         155
Ala Ala Phe Phe Gln Asp Cys Ile Ser Glu Gln Asp Leu Asp Glu Met
Asn Ile Glu Ile Ile Arg Asn Thr Leu Tyr Lys Ala Tyr Leu Glu Ser
                                 185
 Phe Tyr Lys Phe Cys Thr Leu Leu Gly Gly Thr Thr Ala Asp Ala Met
 Cys Pro Ile Leu Glu Phe Xaa Xaa Gln Thr Val Pro Ser Ser Phe His
Thr Val Xaa Gly Ser Thr Leu Arg Ala Trp Arg Xaa Gly Ser Gly
                                          235
                     230
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<sup>&</sup>lt;210> 47

<sup>&</sup>lt;211> 219

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo sapiens

<sup>&</sup>lt;220>

<sup>&</sup>lt;221> SITE

<sup>&</sup>lt;222> (153)

<223> Xaa equals any of the naturally occurring L-amino acids <400> 47 Leu Ala Ala Pro Asp Leu Ser Lys Pro Arg Gly Tyr His Trp Asp Thr  $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$ Ser Asp Trp Met Pro Ser Val Pro Leu Pro Asp Ile Gln Glu Phe Pro Asn Tyr Glu Val Ile Asp Glu Gln Thr Pro Leu Tyr Ser Ala Asp Pro Asn Ala Ile Asp Thr Asp Tyr Tyr Pro Gly Gly Tyr Asp Ile Glu Ser Asp Phe Pro Pro Pro Pro Glu Asp Phe Pro Ala Ala Asp Glu Leu Pro 65 70 75 80 Pro Leu Pro Pro Glu Phe Ser Asn Gln Phe Glu Ser Ile His Pro Pro Arg Asp Met Pro Ala Ala Gly Ser Leu Gly Ser Ser Ser Arg Asn Arg 105 Gln Arg Phe Asn Leu Asn Gln Tyr Leu Pro Asn Phe Tyr Pro Leu Asp 115 120 125 Met Ser Glu Pro Gln Thr Lys Gly Thr Gly Glu Asn Ser Thr Cys Arg Glu Pro His Ala Pro Tyr Pro Pro Kaa Tyr Gln Arg His Phe Glu Ala Pro Ala Val Glu Ser Met Pro Met Ser Val Tyr Ala Ser Thr Ala Ser Cys Ser Asp Val Ser Ala Cys Cys Glu Val Glu Ser Glu Val Met Met 185 Ser Asp Tyr Glu Ser Gly Asp Asp Gly His Phe Glu Glu Val Thr Ile Pro Pro Leu Asp Ser Gln Gln His Thr Glu Val

<210> 48

<211> 49

<212> PRT

<213> Homo sapiens

<400> 48

Met Gly His Cys Cys Leu His Cys Phe Pro His Leu His Gly His Arg
1 5 10 15

Ala Ala Pro Ala Ala Ala Gly Arg Pro Leu Leu Leu Gln Leu 20 25 30

Pro Arg Ala Pro Gln Gly Lys Pro Gln Glu Gly Lys Thr Gln Gly Ser 35 40 45

Gly

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<210> 49
      <211> 26
      <212> PRT
      <213> Homo sapiens
      <400> 49
      Cys Val Ser Leu Thr Ser Ser Thr Gly Phe Gln Lys Trp Lys Asp Val
                                             10
      Pro Cys Glu Asp Lys Phe Ser Phe Val Cys
      <210> 50
      <211> 22
      <212> PRT
      <213> Homo sapiens
      <400> 50
      Ile Met Lys Asp Leu Asp Gln Cys Arg Asp Gly Lys Val Gly Phe Gln
1 5 10 15
13
Ser Phe Phe Ser Leu Ile
4.10
                   20
L.
       <210> 51
       <211> 18
200
       <212> PRT
ļ.
       <213> Homo sapiens
į.
<400> 51
       Gly Lys Tyr Gln Leu Gln Ser Gln Glu Asn Phe Glu Ala Phe Met Lys
                                              10
ļ.i
       Ala Ile
       <210> 52
       <211> 9
       <212> PRT
       <213> Homo sapiens
       <400> 52
       Ile Ala Thr Tyr Arg Ser Leu Leu Glu
       <210> 53
        <211> 12
        <212> PRT
       <213> Homo sapiens
        <400> 53
       Asp Ser Cys Gln Gly Asp Ser Gly Gly Pro Leu Val
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<210> 54

Ala Arg Ala Pro Pro Ala Leu Leu Lys Ile Arg Ser Lys Glu Gly Arg

Cys Ala Gln Pro Ser Arg Thr Ile Gln Thr Ile Cys Leu Pro Ser Met

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Tyr Asn Asp Pro Gln Phe Gly Thr Ser Cys Glu Ile Thr Gly Phe Gly
       Lys Glu Asn Ser Ser Lys
       <210> 55
       <211> 8
       <212> PRT
¥II
       <213> Homo sapiens
Į.
       <400> 55
       Gly Lys Pro Gln Glu Gly Lys Thr
۳.,
IJ
       <210> 56
       <211> 8
į.
       <212> PRT
<213> Homo sapiens
i.i.
Taring
Grand
       <400> 56
       Leu Gly Phe Pro Cys Asn Gln Phe 1
```

<211> 54 <212> PRT

<400> 54

<213> Homo sapiens